

#4

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/904,987

DATE: 10/18/2001
TIME: 09:47:02

Input Set : A:\ES.txt
Output Set: N:\CRF3\10182001\I904987.raw

3 <110> APPLICANT: Novactyl, Inc.
5 <120> TITLE OF INVENTION: Methods and Compositions for Controlling Pathological and
Prepathological
6 Protein Assembly or Aggregation
8 <130> FILE REFERENCE: 42108/26146
10 <140> CURRENT APPLICATION NUMBER: 09/904,987
11 <141> CURRENT FILING DATE: 2001-07-12
13 <160> NUMBER OF SEQ ID NOS: 7
15 <170> SOFTWARE: PatentIn version 3.0
17 <210> SEQ ID NO: 1
18 <211> LENGTH: 43
19 <212> TYPE: PRT
20 <213> ORGANISM: homo sapiens
22 <300> PUBLICATION INFORMATION:
23 <308> DATABASE ACCESSION NO: NCBI ENTREZ / QRHUA4
24 <309> DATABASE ENTRY DATE: 2000-09-15
25 <313> RELEVANT RESIDUES: (672)..(714)
27 <400> SEQUENCE: 1
29 Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys
30 1 5 10 15
32 Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile
33 20 25 30
35 Gly Leu Met Val Gly Val Val Ile Ala Thr
36 35 40
38 <210> SEQ ID NO: 2
39 <211> LENGTH: 770
40 <212> TYPE: PRT
41 <213> ORGANISM: homo sapiens
43 <300> PUBLICATION INFORMATION:
44 <308> DATABASE ACCESSION NO: NCBI ENTREZ / QRHUA4
45 <309> DATABASE ENTRY DATE: 2000-09-15
46 <313> RELEVANT RESIDUES: (1)..(770)
48 <400> SEQUENCE: 2
50 Met Leu Pro Gly Leu Ala Leu Leu Leu Ala Ala Trp Thr Ala Arg
51 1 5 10 15
53 Ala Leu Glu Val Pro Thr Asp Gly Asn Ala Gly Leu Leu Ala Glu Pro
54 20 25 30
56 Gln Ile Ala Met Phe Cys Gly Arg Leu Asn Met His Met Asn Val Gln
57 35 40 45
59 Asn Gly Lys Trp Asp Ser Asp Pro Ser Gly Thr Lys Thr Cys Ile Asp
60 50 55 60
62 Thr Lys Glu Gly Ile Leu Gln Tyr Cys Gln Glu Val Tyr Pro Glu Leu
63 65 70 75 80
65 Gln Ile Thr Asn Val Val Glu Ala Asn Gln Pro Val Thr Ile Gln Asn
66 85 90 95
68 Trp Cys Lys Arg Gly Arg Lys Gln Cys Lys Thr His Pro His Phe Val
69 100 105 110
71 Ile Pro Tyr Arg Cys Leu Val Gly Glu Phe Val Ser Asp Ala Leu Leu

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72	115	120	125
74	Val Pro Asp Lys Cys Lys Phe Leu His Gln Glu Arg Met Asp Val Cys		
75	130	135	140
77	Glu Thr His Leu His Trp His Thr Val Ala Lys Glu Thr Cys Ser Glu		
78	145	150	155
80	Lys Ser Thr Asn Leu His Asp Tyr Gly Met Leu Leu Pro Cys Gly Ile		160
81	165	170	175
83	Asp Lys Phe Arg Gly Val Glu Phe Val Cys Cys Pro Leu Ala Glu Glu		
84	180	185	190
86	Ser Asp Asn Val Asp Ser Ala Asp Ala Glu Glu Asp Asp Ser Asp Val		
87	195	200	205
89	Trp Trp Gly Gly Ala Asp Thr Asp Tyr Ala Asp Gly Ser Glu Asp Lys		
90	210	215	220
92	Val Val Glu Val Ala Glu Glu Glu Val Ala Glu Val Glu Glu Glu		240
93	225	230	235
95	Glu Ala Asp Asp Asp Glu Asp Asp Gly Asp Glu Val Glu Glu		255
96	245	250	
98	Glu Ala Glu Glu Pro Tyr Glu Glu Ala Thr Glu Arg Thr Thr Ser Ile		
99	260	265	270
101	Ala Thr Thr Thr Thr Thr Glu Ser Val Glu Val Val Arg		
102	275	280	285
104	Glu Val Cys Ser Glu Gln Ala Glu Thr Gly Pro Cys Arg Ala Met Ile		
105	290	295	300
107	Ser Arg Trp Tyr Phe Asp Val Thr Glu Gly Lys Cys Ala Pro Phe Phe		320
108	305	310	315
110	Tyr Gly Gly Cys Gly Gly Asn Arg Asn Asn Phe Asp Thr Glu Glu Tyr		335
111	325	330	
113	Cys Met Ala Val Cys Gly Ser Ala Met Ser Gln Ser Leu Leu Lys Thr		
114	340	345	350
116	Thr Gln Glu Pro Leu Ala Arg Asp Pro Val Lys Leu Pro Thr Thr Ala		
117	355	360	365
119	Ala Ser Thr Pro Asp Ala Val Asp Lys Tyr Leu Glu Thr Pro Gly Asp		
120	370	375	380
122	Glu Asn Glu His Ala His Phe Gln Lys Ala Lys Glu Arg Leu Glu Ala		
123	385	390	395
125	Lys His Arg Glu Arg Met Ser Gln Val Met Arg Glu Trp Glu Glu Ala		400
126	405	410	415
128	Glu Arg Gln Ala Lys Asn Leu Pro Lys Ala Asp Lys Lys Ala Val Ile		
129	420	425	430
131	Gln His Phe Gln Glu Lys Val Glu Ser Leu Glu Gln Glu Ala Ala Asn		
132	435	440	445
134	Glu Arg Gln Gln Leu Val Glu Thr His Met Ala Arg Val Glu Ala Met		
135	450	455	460
137	Leu Asn Asp Arg Arg Arg Leu Ala Leu Glu Asn Tyr Ile Thr Ala Leu		
138	465	470	475
140	Gln Ala Val Pro Pro Arg Pro Arg His Val Phe Asn Met Leu Lys Lys		480
141	485	490	495
143	Tyr Val Arg Ala Glu Gln Lys Asp Arg Gln His Thr Leu Lys His Phe		
144	500	505	510

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146 Glu His Val Arg Met Val Asp Pro Lys Lys Ala Ala Gln Ile Arg Ser
147 515 520 525
149 Gln Val Met Thr His Leu Arg Val Ile Tyr Glu Arg Met Asn Gln Ser
150 530 535 540
152 Leu Ser Leu Leu Tyr Asn Val Pro Ala Val Ala Glu Glu Ile Gln Asp
153 545 550 555 560
155 Glu Val Asp Glu Leu Leu Gln Lys Glu Gln Asn Tyr Ser Asp Asp Val
156 565 570 575
158 Leu Ala Asn Met Ile Ser Glu Pro Arg Ile Ser Tyr Gly Asn Asp Ala
159 580 585 590
161 Leu Met Pro Ser Leu Thr Glu Thr Lys Thr Val Glu Leu Leu Pro
162 595 600 605
164 Val Asn Gly Glu Phe Ser Leu Asp Asp Leu Gln Pro Trp His Ser Phe
165 610 615 620
167 Gly Ala Asp Ser Val Pro Ala Asn Thr Glu Asn Glu Val Glu Pro Val
168 625 630 635 640
170 Asp Ala Arg Pro Ala Ala Asp Arg Gly Leu Thr Thr Arg Pro Gly Ser
171 645 650 655
173 Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser Glu Val Lys Met Asp
174 660 665 670
176 Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys Leu
177 675 680 685
179 Val Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile Gly
180 690 695 700
182 Leu Met Val Gly Val Val Ile Ala Thr Val Ile Val Ile Thr Leu
183 705 710 715 720
185 Val Met Leu Lys Lys Gln Tyr Thr Ser Ile His His Gly Val Val
186 725 730 735
188 Glu Val Asp Ala Ala Val Thr Pro Glu Glu Arg His Leu Ser Lys Met
189 740 745 750
191 Gln Gln Asn Gly Tyr Glu Asn Pro Thr Tyr Lys Phe Phe Glu Gln Met
192 755 760 765
194 Gln Asn
195 770
197 <210> SEQ ID NO: 3
198 <211> LENGTH: 253
199 <212> TYPE: PRT
200 <213> ORGANISM: homo sapiens
202 <300> PUBLICATION INFORMATION:
203 <308> DATABASE ACCESSION NO: NCBI ENTREZ / XM_009567
204 <309> DATABASE ENTRY DATE: 2001-04-17
205 <313> RELEVANT RESIDUES: (1)...(253)
207 <400> SEQUENCE: 3
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210 1 5 10 15
212 Ser Asp Leu Gly Leu Cys Lys Lys Arg Pro Lys Pro Gly Gly Trp Asn
213 20 25 30
215 Thr Gly Gly Ser Arg Tyr Pro Gly Gln Gly Ser Pro Gly Gly Asn Arg
216 35 40 45

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218 Tyr Pro Pro Gln Gly Gly Gly Trp Gly Gln Pro His Gly Gly Gly
219 50 55 60
221 Trp Gly Gln Pro His Gly Gly Gly Trp Gly Gln Pro His Gly Gly Gly
222 65 70 75 80
224 Trp Gly Gln Pro His Gly Gly Gly Trp Gly Gln Gly Gly Thr His
225 85 90 95
227 Ser Gln Trp Asn Lys Pro Ser Lys Pro Lys Thr Asn Met Lys His Met
228 100 105 110
230 Ala Gly Ala Ala Ala Gly Ala Val Val Gly Gly Leu Gly Gly Tyr
231 115 120 125
233 Met Leu Gly Ser Ala Met Ser Arg Pro Ile Ile His Phe Gly Ser Asp
234 130 135 140
236 Tyr Glu Asp Arg Tyr Tyr Arg Glu Asn Met His Arg Tyr Pro Asn Gln
237 145 150 155 160
239 Val Tyr Tyr Arg Pro Met Asp Glu Tyr Ser Asn Gln Asn Asn Phe Val
240 165 170 175
242 His Asp Cys Val Asn Ile Thr Ile Lys Gln His Thr Val Thr Thr
243 180 185 190
245 Thr Lys Gly Glu Asn Phe Thr Glu Thr Asp Val Lys Met Met Glu Arg
246 195 200 205
248 Val Val Glu Gln Met Cys Ile Thr Gln Tyr Glu Arg Glu Ser Gln Ala
249 210 215 220
251 Tyr Tyr Gln Arg Gly Ser Ser Met Val Leu Phe Ser Ser Pro Pro Val
252 225 230 235 240
254 Ile Leu Leu Ile Ser Phe Leu Ile Phe Leu Ile Val Gly
255 245 250
257 <210> SEQ ID NO: 4
258 <211> LENGTH: 140
259 <212> TYPE: PRT
260 <213> ORGANISM: homo sapiens
262 <300> PUBLICATION INFORMATION:
263 <308> DATABASE ACCESSION NO: NCBI ENTREZ / XM_003494
264 <309> DATABASE ENTRY DATE: 2001-04-16
265 <313> RELEVANT RESIDUES: (1)..(140)
267 <400> SEQUENCE: 4
269 Met Asp Val Phe Met Lys Gly Leu Ser Lys Ala Lys Glu Gly Val Val
270 1 5 10 15
272 Ala Ala Ala Glu Lys Thr Lys Gln Gly Val Ala Glu Ala Ala Gly Lys
273 20 25 30
275 Thr Lys Glu Gly Val Leu Tyr Val Gly Ser Lys Thr Lys Glu Gly Val
276 35 40 45
278 Val His Gly Val Ala Thr Val Ala Glu Lys Thr Lys Glu Gln Val Thr
279 50 55 60
281 Asn Val Gly Gly Ala Val Val Thr Gly Val Thr Ala Val Ala Gln Lys
282 65 70 75 80
284 Thr Val Glu Gly Ala Gly Ser Ile Ala Ala Ala Thr Gly Phe Val Lys
285 85 90 95
287 Lys Asp Gln Leu Gly Lys Asn Glu Glu Gly Ala Pro Gln Glu Gly Ile
288 100 105 110

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290 Leu Glu Asp Met Pro Val Asp Pro Asp Asn Glu Ala Tyr Glu Met Pro
291 115 120 125
293 Ser Glu Glu Gly Tyr Gln Asp Tyr Glu Pro Glu Ala
294 130 135 140
296 <210> SEQ ID NO: 5
297 <211> LENGTH: 758
298 <212> TYPE: PRT
299 <213> ORGANISM: homo sapiens
301 <300> PUBLICATION INFORMATION:
302 <308> DATABASE ACCESSION NO: NCBI ENTREZ / NM_016835
303 <309> DATABASE ENTRY DATE: 2001-02-13
304 <313> RELEVANT RESIDUES: (1)..(758)
306 <400> SEQUENCE: 5
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311 Thr Tyr Gly Leu Gly Asp Arg Lys Asp Gln Gly Gly Tyr Thr Met His
312 20 25 30
314 Gln Asp Gln Glu Gly Asp Thr Asp Ala Gly Leu Lys Glu Ser Pro Leu
315 35 40 45
317 Gln Thr Pro Thr Glu Asp Gly Ser Glu Glu Pro Gly Ser Glu Thr Ser
318 50 55 60
320 Asp Ala Lys Ser Thr Pro Thr Ala Glu Asp Val Thr Ala Pro Leu Val
321 65 70 75 80
323 Asp Glu Gly Ala Pro Gly Lys Gln Ala Ala Ala Gln Pro His Thr Glu
324 85 90 95
326 Ile Pro Glu Gly Thr Thr Ala Glu Glu Ala Gly Ile Gly Asp Thr Pro
327 100 105 110
329 Ser Leu Glu Asp Glu Ala Ala Gly His Val Thr Gln Glu Pro Glu Ser
330 115 120 125
332 Gly Lys Val Val Gln Glu Gly Phe Leu Arg Glu Pro Gly Pro Pro Gly
333 130 135 140
335 Leu Ser His Gln Leu Met Ser Gly Met Pro Gly Ala Pro Leu Leu Pro
336 145 150 155 160
338 Glu Gly Pro Arg Glu Ala Thr Arg Gln Pro Ser Gly Thr Gly Pro Glu
339 165 170 175
341 Asp Thr Glu Gly Gly Arg His Ala Pro Glu Leu Leu Lys His Gln Leu
342 180 185 190
344 Leu Gly Asp Leu His Gln Glu Gly Pro Pro Leu Lys Gly Ala Gly Gly
345 195 200 205
347 Lys Glu Arg Pro Gly Ser Lys Glu Glu Val Asp Glu Asp Arg Asp Val
348 210 215 220
350 Asp Glu Ser Ser Pro Gln Asp Ser Pro Pro Ser Lys Ala Ser Pro Ala
351 225 230 235 240
353 Gln Asp Gly Arg Pro Pro Gln Thr Ala Ala Arg Glu Ala Thr Ser Ile
354 245 250 255
356 Pro Gly Phe Pro Ala Glu Gly Ala Ile Pro Leu Pro Val Asp Phe Leu
357 260 265 270
359 Ser Lys Val Ser Thr Glu Ile Pro Ala Ser Glu Pro Asp Gly Pro Ser
360 275 280 285

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/904,987

DATE: 10/18/2001
TIME: 09:47:03

Input Set : A:\ES.txt
Output Set: N:\CRF3\10182001\I904987.raw